

SEQUENCE LISTING

<110> Perez, Pascual
Gutierrez-Marcos, Jose
Dickinson, Hugh

<120> MEG1 Endosperm-Specific Promoter and Genes

<130> 11887*8

<150> PCT/EP04/052760
<151> 2004-11-04

<150> EP 03292739.4
<151> 2003-11-03

<160> 64

<170> PatentIn version 3.2

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atagataact tacaattttg totaaaagag actaaatcac tgctaagttt ggtctttggt 240
gaatacttgc cagtgaattg gttttcgcta tagtatatat ataagtatac actcttctag 300
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cgctatagta	tatatatata	agtatacact	cttctaggat	tatagtatat	atatatatat		360
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 35 40 45
 Asn Gly Ala Arg Cys Val Val Gly Phe Pro Pro Cys Lys Asp Asn Lys
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gataacaagt gctactgctg cattgggggg cgaactcatg ctcgctactc tacgctggct      660
gagtgtagtc atgcctgctt ctaaacaaaa attaagatca ctggtattat atacattgta      720
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Cys Cys Ile Gly Gly Asp Val Gly Phe Pro Pro Cys Lys Asp Asn Lys
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Cys Tyr Cys Cys Ile Gly Gly Arg Thr His Ala Arg Tyr Ser Thr Leu
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Ala Glu Cys Ser His Ala Cys Phe
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20 25 30

Ile Met Gln Gly Asn Gly Ala Arg Cys Val Val Gly Phe Pro Pro Cys
 35 40 45

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Tyr Ser Thr Met Ala Glu Cys Ser His Ala Cys Phe
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 35 40 45
 Val Asn Lys Glu Ser Arg His Pro Gly Gly Asp Pro His Val Leu Cys
 50 55 60
 Phe Val Asp Phe Asp Asn Pro Ala Gln Ala Thr Ile Ala Leu Glu Ala
 65 70 75 80
 Leu Gln Gly His Val Thr Asp Asp Val Asn Val Ser Ala Pro Ala Glu
 85 90 95
 Glu Gly Ile Leu Arg Glu Lys Arg Ala Gln Cys Ala Gln Gly Phe Leu
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ataacttttg gatgccaacg gaggtcgttc caccogtgga caacatggac atgatgcaag	240
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Met Asn Ser Leu Ile Thr Ser His Ile Ala Asn Asn Thr Tyr Thr Asn
          20          25          30

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Asn Asn Gln His Val Val Ala Ser Arg Ser Ala Ile Val Asn His Asn
          35          40          45

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Asn Phe Gly Met Pro Thr Glu Val Val Pro Pro Val Asp Asn Met Asp
          50          55          60

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Met Met Gln Gly Tyr Leu Met Ala Asp Thr Asp Ala Met Arg Leu Val
65          70          75          80

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Gln Gly Gln Gln His Met Pro Asn Val Val Pro Asn Gln Arg Arg His

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85								90				95			
Ala	Val	Lys	Phe	Trp	Thr	Thr	Asp	Glu	His	Arg	Asn	Phe	Leu	Arg	Gly
			100					105					110		
Leu	Glu	Val	Phe	Gly	Arg	Gly	Lys	Trp	Lys	Asn	Ile	Ser	Lys	Tyr	Phe
		115					120					125			
Val	Pro	Thr	Arg	Thr	Pro	Val	Gln	Ile	Ser	Ser	His	Ala	Gln	Lys	Tyr
	130					135					140				
Phe	Arg	Arg	Gln	Glu	Cys	Thr	Thr	Glu	Lys	Gln	Arg	Phe	Ser	Ile	Asn
145					150					155					160
Asp	Val	Gly	Leu	Tyr	Asp	Thr	Gln	Pro	Trp	Val	Arg	Gln	Asn	Asn	Ser
			165						170					175	
Ser	Ser	Ser	Trp	Glu	Ala	Leu	Thr	Phe	Thr	Ala	Gly	Arg	Ala	Tyr	Asn
			180					185					190		
Asn	Thr	Asn	Tyr	Cys	Ala	Phe	Asn	Ser	Leu	Pro	Tyr	Ala	Ser	Ser	Gln
	195						200					205			
Ala	Ser	Asn	Asn	Gln	Val	Ala	Thr	Trp	Ile	Thr	Asp	Gln	Gln	Ala	Thr
	210					215					220				
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225					230					235					240
Asn Arg															

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 <211> 83
 <212> PRT
 <213> Zea mays

<220>
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 <223> MEG1-3, second ORF

<400> 53

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			20					25					30		

Ser Thr Pro Ala Lys Glu Gly Ile Met Gln Gly Asn Gly Ala Arg Cys
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Asp Val Gly Phe Pro Pro Cys Lys Asp Asn Lys Cys Tyr Cys Cys Ile
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Gly Gly Arg Thr His Ala Arg Tyr Ser Thr Leu Ala Glu Cys Ser His
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Ala Cys Phe

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<400> 56
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aattctgtgt actttgattt atattatgta aattactcta gtcttttata ttatttctta 180
ctctttattg ttattcgaag cattgtgtta tgatgagtca tttatgtaat tgctatgtac 240
gtgagttttg atcctagcac gtacatgggt cgcattcggg ttaccttcta aaacctgggg 300
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ggtagtagta gagccagaat tgtaaccttg ggttttccca cacctcaa atagatagat 480
atagggatat agatagatat agcaaattca ccaaataata taggggtata gatatagata 540
taagaagggg tatagatata gatatagata tatagaagat atagatagat agatagatat 600
gatagaatag ataacttaca attttgtcta aaagaaacta aatcactgct aagtttggag 660
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agtatacact cttctaggat tatagtatat atatatatat aagtatacac tcttctagga 780
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cgctagtgtt tttctcgtta cttctcctcg gatactttgc tgctcatgca catgggaagg 900
gtaagtga aaactatacaga catgtgtgtg catgcttaga tagatctaga caatttagaa 960
gatgttatta tatgataccg tgtgtatcat ggcgaattgc taatgtatcg caatcccctg 1020
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gctaaagaag gaattatgca aggaacgga gcacgatgcg ttgtagggtt tcctccatgc 1200
aaagataaca agtgctactg ctgcattggg gggcgaaactc atgctcgcta ctctcgatgg 1260
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atttataata ttatggaatt agttgtatat 1350

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 <223> nucleotides 1-127 of promoter MEG1-1

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<210> 60
 <211> 25
 <212> DNA
 <213> Artificial

<220>
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<400> 60
 acacctcaaa tagatatgga tatag 25

<210> 61
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<220>
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<400> 61
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 agaattgtaa ccttgggttt tcccacacct caaatagata tagatatagg gatatagata 180

gatatagcaa attcaccaaa taatataggg gtatagatat agatataaga aggggtatag	240
atatagatat agatatatag aagatataga tagatagata gatatgatag aatagataac	300
ttacaatttt gtctaaaaga aactaaatca ctgctaagtt tggagtagca tatctttggt	360
gaatacttgc tagtgaattg gtttccgcta tagtatatat atataagtat acactcttct	420
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<220>
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<400> 63	
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<210> 64
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<220>
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